

0100
23-01.. 0280

OIPE

ENTERED

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/817,360

DATE: 04/05/2001

TIME: 12:14:04

Input Set : A:\SeqList for UCSF-129CIP.txt

Output Set: N:\CRF3\04052001\I817360.raw

4 <110> APPLICANT: German, Michael S.
5 Lin, Joseph
7 <120> TITLE OF INVENTION: PRODUCTION OF PANCREATIC ISLET CELLS
8 AND DELIVERY OF INSULIN
13 <130> FILE REFERENCE: UCSF-129CIP
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/817,360
C--> 15 <141> CURRENT FILING DATE: 2001-03-20
15 <150> PRIOR APPLICATION NUMBER: 09/535,145
16 <151> PRIOR FILING DATE: 2000-03-24
18 <150> PRIOR APPLICATION NUMBER: 60/128,180
19 <151> PRIOR FILING DATE: 1999-04-06
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23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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27 <212> TYPE: DNA
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33 tctaacttgc agttaataa atcaggcaag gctggcctat gaggcagaca agtgtgaaga 180
34 aggagaagga ggaggagaag gagaaggaga aagaagaaga aggaggagaa gaagaagaag 240
35 aagaagaaga agaaggaggag gaggaggagg aggaggagga agcagcagca gcagcagcag 300
36 cttgaatgga cagtggttcc ccttgccctag aaaatgggac cattatttct tttctaattct 360
37 gacccccaga ctccaggactt cctctatttt ctgcattttg ggtctctctg ttttgccttg 420
38 aaaaaaaatg ttttctccca aatcaaggag cagtagctgg tgcaaggga aatctagggc 480
39 taggagtctt aagatatgac ttctatgtgg ttctgataga acttgctggg tgaccttgag 540
40 agagtcactc cccctctctg ggccttgatt ttttcatott taaagaaggc ctcaaattcc 600
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42 aaaatgctaa cctgacagtc ccagatggtc cctttattgg ttctgacctt ggtctcaggc 720
43 ttcattttcc cacagcaagg gaaggagcct gctcacagag caccagctaa gatcagcagg 780
44 accgcgccac acccccgcgc agtccctagag ccccccctct gctggttcct gagcatacca 840
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46 gggcagactc acagacctac ctcatccctc caccacccc tacagtctcg aagtcgggtc 960
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58 gtctcagcta ctgggaaggc tgaagtggga ggatagcttg agcccaggag atcaaggctg 1680
59 cagtgaagctg tgattgcacc actgcagtc agcctggggc acagaaggag accgtttttt 1740

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62	gggatactat	ctacaagagt	tagtaggctt	gtagtaatgg	aatgtccgct	tgaggggtcc	1920
63	ccgcacagcc	aacccccggc	tctggagtg	gatctatggg	ggtgggggtc	taagcgctc	1980
64	tggggagtgt	gaggtagcat	ctcaggggtg	ggcagaggct	cggacacccc	caaaaggtct	2040
65	gtgaatggaa	gggacatagg	caggatctct	ctcagtgtg	tcccctgtct	tccaggatga	2100
66	agagaggcag	tgaaacacca	ggagagcagg	gcgtccttta	gaattcctgg	acccttctcc	2160
67	aggctgctag	tcaggacaat	gagctcgtgg	ttgtctttgc	cactatcttc	ctgtgcgatt	2220
68	tcagacaagc	cacctccctc	actaagccta	aatttcccca	tgtgtaacgt	gcaggcattg	2280
69	taccgtāgag	gcatacaagt	cccctccagg	acagatgcta	aggaaagata	ggctaggagc	2340
70	aaagccgtct	gaggtggcct	gaccagagcc	acacgaggct	cttctcactg	ggcgaggctc	2400
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102	accaggctgt	gaactgggaa	agcgctagcc	cggccaggga	gcatctcccc	agcctccctg	4320
103	cgaactgcgc	ctgaaacgtg	agctgcgctg	caggtgcctg	gagcacccgc	catctttttt	4380
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106	aaagcctata	ttttggatat	tagattattt	agggattgct	ggatgacatt	taagccaata	4560
107	aaaaaaaaatg	gaccttcaag	aagccttggc	aagatgactc	cattgtgtgt	tggggagagg	4620
108	agggccacag	tcactacagc	tgaggaagag	cacttctgtc	caaagagagg	gatgacactc	4680

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Input Set : A:\SeqList for UCSF-129CIP.txt

Output Set : N:\CRF3\04052001\I817360.raw

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109 tttctggagg tctgggctag agccagggca gattggggtt ggagagctgg aagtcttcta 4740
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111 gagtgggtta gaagaaagac aagcctccaa ctaggacaac tgactctcac ttgctggccc 4860
112 tttccccaac tccaccagcc tagctttaga gcaactgttg gttgcacttg gggaaaggat 4920
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116 gagcgtctgg ctggaatata cagcaacgcc ccctccctca tcacctggca gccttgattg 5160
117 aaaacttatt aagaactgt tcaaggtttc cagccacacc atgtctctta ctggcaagggt 5220
118 ggaataggac tgggtcagca tgagcactga aatctgtccc aggagtgccca gtagagcacc 5280
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120 <210> SEQ ID NO: 2

121 <211> LENGTH: 214

122 <212> TYPE: PRT

123 <213> ORGANISM: Homo Sapiens

125 <400> SEQUENCE: 2

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129 20 25 30
130 Thr Ser Ala Pro Pro Ser Pro Thr Arg Thr Arg Gly Asn Cys Ala Glu
131 35 40 45
132 Ala Glu Glu Gly Gly Cys Arg Gly Ala Pro Arg Lys Leu Arg Ala Arg
133 50 55 60
134 Arg Gly Gly Arg Ser Arg Pro Lys Ser Glu Leu Ala Leu Ser Lys Gln
135 65 70 75 80
136 Arg Arg Ser Arg Arg Lys Lys Ala Asn Asp Arg Glu Arg Asn Arg Met
137 85 90 95
138 His Asn Leu Asn Ser Ala Leu Asp Ala Leu Arg Gly Val Leu Pro Thr
139 100 105 110
140 Phe Pro Asp Asp Ala Lys Leu Thr Lys Ile Glu Thr Leu Arg Phe Ala
141 115 120 125
142 His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
143 130 135 140
144 Ser Leu Tyr Ala Leu Glu Pro Pro Ala Pro His Cys Gly Glu Leu Gly
145 145 150 155 160
146 Ser Pro Gly Gly Ser Pro Gly Asp Trp Gly Ser Leu Tyr Ser Pro Val
147 165 170 175
148 Ser Gln Ala Gly Ser Leu Ser Pro Ala Ala Ser Leu Glu Glu Arg Pro
149 180 185 190
150 Gly Leu Leu Gly Ala Thr Ser Ser Ala Cys Leu Ser Pro Gly Ser Leu
151 195 200 205
152 Ala Phe Ser Asp Phe Leu
153 210
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155 <210> SEQ ID NO: 3

156 <211> LENGTH: 1861

157 <212> TYPE: DNA

158 <213> ORGANISM: Mus musculus

160 <400> SEQUENCE: 3

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164 ctaatgggag ctcctcccag ctgccagcca agaagacact tgactccttg atcgctggtt 240
165 catttagaca agccgtttcc ctctctgagc caaaagaccc catgtgtaat actcaaagaa 300
166 gaggccttcc ttatatatat ataggcacc ccaaacctcc ttcattgctac caagaaaggg 360
167 tctggacaca tgccaaaaag aaagaggaaa aggcaaaagt ctcccagcg gccggacggg 420
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169 cgggggaggc gtggagtgagg ggaacaaaca gagtgtgtgt cccctcccc gaccctgcc 540
170 ctctgtccgg aactccagctg tgctctgcgg gtgggggagg tggggggagg agcgggctcg 600
171 cgtggcgagc cccctgggccc cccctcgctg attggcccg gtgacaggca gcagcccgcc 660
172 aggcacgctc ctggccgggg gcagagcaga taaagcgtgc caggggacac acgacttgca 720
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190 gacctgtctg gctcagggtg gtgggtgcta ttggaaagg aggggaccag agccgtctgg 1800
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192 c 1861
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194 <210> SEQ ID NO: 4

195 <211> LENGTH: 214

196 <212> TYPE: PRT

197 <213> ORGANISM: Mus musculus

199 <400> SEQUENCE: 4

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202 Thr Gln Gln Pro Phe Pro Gly Ala Ser Asp His Glu Val Leu Ser Ser
203 20 25 30
204 Asn Ser Thr Pro Pro Ser Pro Thr Leu Ile Pro Arg Asp Cys Ser Glu
205 35 40 45
206 Ala Glu Val Gly Asp Cys Arg Gly Thr Ser Arg Lys Leu Arg Ala Arg
207 50 55 60
208 Arg Gly Gly Arg Asn Arg Pro Lys Ser Glu Leu Ala Leu Ser Lys Gln
209 65 70 75 80
210 Arg Arg Ser Arg Arg Lys Lys Ala Asn Asp Arg Glu Arg Asn Arg Met
211 85 90 95
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216 His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
217          130          135          140
218 Ser Phe Tyr Gly Pro Glu Pro Pro Val Pro Cys Gly Glu Leu Gly Ser
219 145          150          155          160
220 Pro Gly Gly Gly Ser Asn Gly Asp Trp Gly Ser Ile Tyr Ser Pro Val
221          165          170          175
222 Ser Gln Ala Gly Asn Leu Ser Pro Thr Ala Ser Leu Glu Glu Phe Pro
223          180          185          190
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225          195          200          205
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232 <213> ORGANISM: Artificial Sequence
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235 <223> OTHER INFORMATION: oligonucleotide primer
237 <400> SEQUENCE: 5
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240 <210> SEQ ID NO: 6
241 <211> LENGTH: 23
242 <212> TYPE: DNA
243 <213> ORGANISM: Artificial Sequence
245 <220> FEATURE:
246 <223> OTHER INFORMATION: primer
248 <400> SEQUENCE: 6
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251 <210> SEQ ID NO: 7
252 <211> LENGTH: 16
253 <212> TYPE: DNA
254 <213> ORGANISM: Artificial Sequence
256 <220> FEATURE:
257 <223> OTHER INFORMATION: oligonucleotide primer
259 <400> SEQUENCE: 7
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263 <211> LENGTH: 21
264 <212> TYPE: DNA
265 <213> ORGANISM: Artificial Sequence
267 <220> FEATURE:
268 <223> OTHER INFORMATION: oligonucleotide primer
270 <400> SEQUENCE: 8
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273 <210> SEQ ID NO: 9
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VERIFICATION SUMMARY

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L:15 M:270 C: Current Application Number differs, Replaced Current Application No

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date